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Application Serial Number:	09/786,502
Source:	1600
Date Processed by STIC:	9-9-03

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

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Revised 04/24/2003

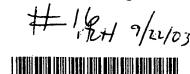




ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: 09/786,502	
ATTN: NEW RULES CASE	s: Please disregard english "	ALPHA" HEADERS WHICH WERE INSERTED BY PTO	SOFTWARE
Wrapped Nucleics Wrapped Aminos		e "wrapped" down to the next line. This may occur if your file r creating it. Please adjust your right margin to .3; this will	
2Invalid Line Length	The rules require that a line not exceed	172 characters in length. This includes white agaces.	
3Misaligned Amino Numbering	The numbering under each 5th amino acuse space characters, instead.	cid is misaligned. Do not use tab codes between numbers;	
4Non-ASCII	The submitted file was not saved in As ensure your subsequent submission is	SCII(DOS) text, as required by the Sequence Rule Please is naved in ASCII text.	
5Variable Length	each n or Xaa can only represent a si	epresenting more than one residue. Per Sequence Rules, Ingle residue. Please present the maximum number of each cate in the <220>-<223> section that some may be missing.	
6Patentln 2.0 "bug"	sequences(s)	sed the <220>-<223> section to be missing from amino acid Patentin would automatically generate this section from the Please manually copy the relevant <220>-<223> section to This applies to the mandatory <220>-<223> sections for	
7Skipped Sequences (OLD RULES)	(2) INFORMATION FOR SEQ ID NO (i) SEQUENCE CHARACTE	nal, please insert the following lines for each skipped sequence: 2:X: (insert SEQ ID NO where "X" is shown) RISTICS: (Do not insert any subheadings under this heading) RID NO:X: (insert SEQ ID NO where "X" is shown)	
	Please also adjust the "(ii) NUMBER C	OF SEQUENCES:" response to include the skipped sequences.	
8Skipped Sequences (NEW RULES)	Sequence(s) missing. If Intent(<210> sequence id number <400> sequence id number 000	ional, please insert the following lines for each skipped sequence.	
Use of n's or Xaa's (NEW RULES)		cted in the Sequence Listing. 20>-<223> is MANDATORY if n's or Xaa's are present. ain location of n or Xaa, and which residue n or Xaa represents.	
0lnvalid <213> Response		ralid <213> responses are: Unknown, Artificial Sequence, or >-<223> section is required when <213> response is Unknown or	r
1Use of <220>	Use of <220> to <223> is MANDATOR "Unknown." Please explain source of g	0> "Feature" and associated numeric identifiers and responses. RY if <213> "Organism" response is "Artificial Sequence" or genetic material in <220> to <223> section. ol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)	
2PatentIn 2.0 "bug"	resulting in missing mandatory numeric	on of Patentin version 2.0. This causes a corrupted file, identifiers and responses (as indicated on raw sequence ger" or any other manual means to copy file to floppy disk.	
3Misuse of n	n can only be used to represent a single	nucleotide in a nucleic acid sequence. N is not used to represent	

AMC/MH - Biotechnology Systems Branch - 08/21/2001







Does Not Comply Corrected Diskette Needed

1600

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/786,502

DATE: 09/09/2003 TIME: 07:00:07

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Output Set: N:\CRF4\09092003\I786502.raw

3 <110> APPLICANT: Sadelain, Michel Bander, Neil H Gong, Michael 7 <120> TITLE OF INVENTION: Fusion Receptors Specific For Prostate-Specific Membrane Antigen and Uses Thereof 10 <130> FILE REFERENCE: MSK.P-040 12 <140> CURRENT APPLICATION NUMBER: 09/786,502 13 <141> CURRENT FILING DATE: 2001-05-18 15 <150> PRIOR APPLICATION NUMBER: PCT/US99/20349 16 <151> PRIOR FILING DATE: 1999-09-03 18 <150> PRIOR APPLICATION NUMBER: US 60/099,138 19 <151> PRIOR FILING DATE: 1998-09-04 21 <160> NUMBER OF SEQ ID NOS: 8 23 <170> SOFTWARE: PatentIn version 3.1 25 <210> SEQ ID NO: 1 26 <211> LENGTH: 22 27 <212> TYPE: DNA 28 <213> ORGANISM: synthetic construct 30 <400> SEQUENCE: 1 22 31 aggtsmarct gcagsagtcw gg 34 <210> SEQ ID NO: 2 35 <211> LENGTH: 34 36 <212> TYPE: DNA 37 <213> ORGANISM: synthetic construct 39 <400> SEQUENCE: 2 34 40 tgaggagacg gtgaccgtgg tcccttggcc ccag 43 <210> SEQ ID NO: 3 44 <211> LENGTH: 24 45 <212> TYPE: DNA 46 <213> ORGANISM: synthetic construct 48 <400> SEQUENCE: 3 24 49 gacattgagc tcacccagtc tcca 52 <210> SEQ ID NO: 4 53 <211> LENGTH: 33 54 <212> TYPE: DNA 55 <213> ORGANISM: synthetic construct 57 <400> SEQUENCE: 33 58 tgcggccgcc cgtttgatct ccagcttggt ccc 61 <210> SEQ ID NO: 5 62 <211> LENGTH: 43 63 <212> TYPE: DNA 64 <213> ORGANISM: (synthetic construct

66 <400> SEQUENCE: 5



PATENT APPLICATION: US/09/786,502

DATE: 09/09/2003

TIME: 07:00:07

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Output Set: N:\CRF4\09092003\I786502.raw

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80	<211> LENGTH: 27	
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RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/09/786,502

DATE: 09/09/2003 TIME: 07:00:08

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Invalid Line Length: The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 7

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/786,502

DATE: 09/09/2003

TIME: 07:00:08

Input Set : N:\efs\09786502\ptoms.txt
Output Set: N:\CRF4\09092003\I786502.raw

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